



# SEQUENCE LISTING

<110> Leproust, Eric M.  
Wolber, Paul K.

<120> METHODS FOR IDENTIFYING SUITABLE NUCLEIC  
ACID PROBE SEQUENCES FOR USE IN NUCLEIC ACID ARRAYS

<130> 10030416-1

<140> 10/698,195

<141> 2003-10-30

<160> 9

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 16

<212> DNA

<213> homo sapiens

<400> 1

atggacttag cattcg

16

<210> 2

<211> 12

<212> DNA

<213> homo sapiens

<400> 2

atggacttag ca

12

<210> 3

<211> 12

<212> DNA

<213> homo sapiens

<400> 3

tggacttagc at

12

<210> 4

<211> 12

<212> DNA

<213> homo sapiens

<400> 4

ggacttagca tt

12

<210> 5

<211> 12

<212> DNA

<213> homo sapiens

<400> 5

gacttagcat tc	12
<210> 6	
<211> 12	
<212> DNA	
<213> homo sapiens	
<400> 6	
acttagcatt cg	12
<210> 7	
<211> 25	
<212> DNA	
<213> homo sapiens	
<400> 7	
atcatcgtag ctggtcagtg tatcc	25
<210> 8	
<211> 25	
<212> DNA	
<213> homo sapiens	
<400> 8	
atcatcgtag ctggtcagtg tatcc	25
<210> 9	
<211> 35	
<212> DNA	
<213> homo sapiens	
<400> 9	
atcatcgtag ctggtcagtg tatccaaaaa aaaaa	35